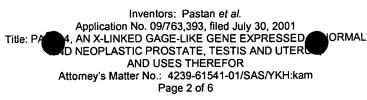
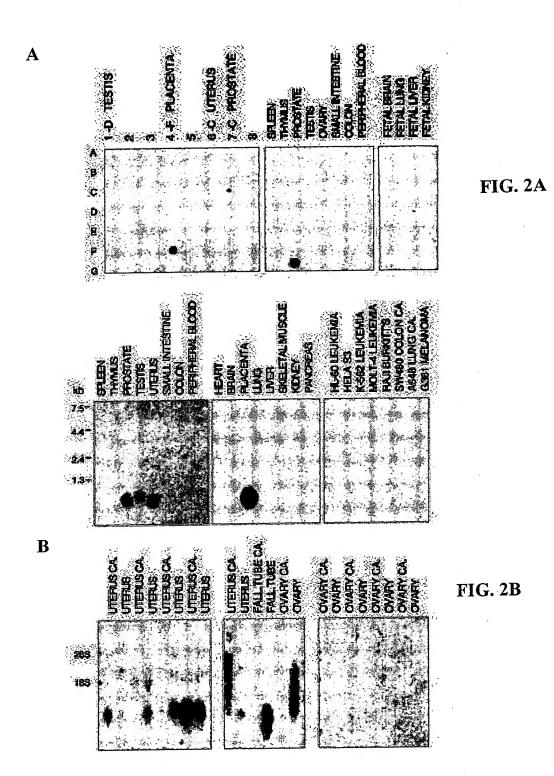
Amendments to the Drawings

The attached sheets of drawings include changes to Figs. 1, 2A, 2B, 2C, 3, 4, and 5. Theses sheets replace the original sheets of drawings.

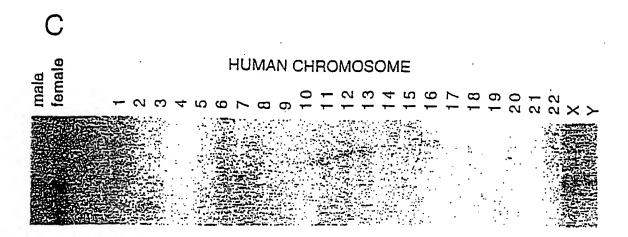
Attachment: Replacement Sheets (6)

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	GAGE1 GAGE3	Y E M m.s K m Q V
•	GAGE4	MA Ö M
	GAGE5,6	m Q V
	2005	: : : : : : MSLEOKSOmEKPEEGLDTOE.EALGLVGVQAATTEEQEAVS
	MAGES MAGES	LG RA QAG. P MD IPA KA
		•
	PAGE	DNQDIEPGQEREGTPPIEERKVEGDCQE
	GAGE2	:: : :
	GAGE1	D
	GAGE3-6	, , , , , , , , , , , , , , , , , , ,
	MAGE5	
	MAGE8	T IM E TDS S PQ. E SSL VT S
	PAGE	MDLEKTRSE <u>RGD</u> GSDVKEKTPPNPKHAKTKEAGDGQP
	a. a	: : : : :
	GAGE2 GAGE1	M . MR HYVA
	GAGE3-5	<u>M</u>
	GAGE6	v
	MAGE5	ROSIKGSSNOEEEGPSTSPDPESVFRAALSKKVADLIHFLLLKY
	MAGE8	S DE SN AHL L E DE E VR R
		•
D		
В	PAGE1	MSARVRSRSRGRGDGQEAPDVVAFVAPGES
	PAGE2: : : . MSELVRARSQSSERGNDQESSQPVGSVIVQEPTEEKR
		MTSF
	PAGE3	
		WE FOOFT
	PAGE1	QEEEPPTDNQGPDMEAFQQEL
	PAGE2	OQEEPPTONODIEPGOEREGTPPIZERKVEGDCQEM
	PAGE3	nktappiesodytpgoerdegaldfovpslaavlwel
		•
	PAGE1	DLEKTRSERGDGSDVKEKTPPNPKHAKTKEAGDGQP
	PAGE2	ALLKIEDEPGDGPDVREGIMPTFDLTKVLEAGDAQP
	-	
	PAGE3	TRPKTGGERGDGPNVKGESLPNLEPVKIPEAGEGQPSV
		•





Inventors: Pastan et al.
Application No. 09/763,393, filed July 30, 2001
AGE-4, AN X-LINKED GAGE-LIKE GENE EXPREST IN NORMAL
AND NEOPLASTIC PROSTATE, TESTIS AND UTERUS,
AND USES THEREFOR
Attorney's Matter No.: 4239-61541-01/SAS/YKH:kam
Page 3 of 6



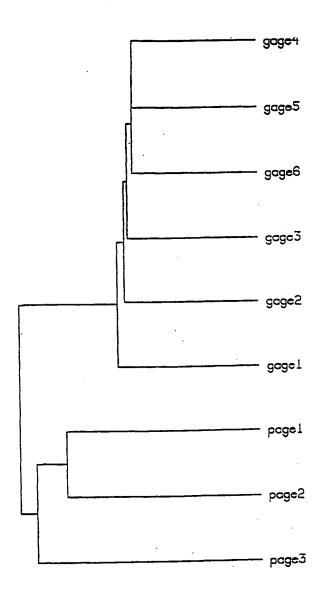


FIG. 4

Inventors: Pastan et al.
Application No. 09/763,393, filed July 30, 2001
Title: PA
AN X-LINKED GAGE-LIKE GENE EXPRESSED
AND NEOPLASTIC PROSTATE, TESTIS AND UTERUS,
AND USES THEREFOR

AND USES THEREFOR Attorney's Matter No.: 4239-61541-01/SAS/YKH:kam Page 5 of 6

liver (cDNA library)
muscle (cDNA library)
neg. control (no DNA)
placenta (cDNA library)
testis tumor (cDNA)
prostate tumor (cDNA)
prostate normal (cDNA)
neg. control (no DNA)
pos. control (plasmid)
molecular weight marker



Attorney's Matter No.: 4239-61541-01/SAS/YKH:kam Page 6 of 6

Page 4 sequence (bold and underline: open reading frame).

GAAGAATTCGCCAGGCTCTCTGCTGACTCAAGTTCTTCAGTTCACGATCTTCTAGTT GCAGCG

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